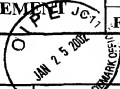


<b>INFORMATION DISCLOSURE STATEMENT</b>	Atty. Docket No.: 6317.N	Serial No.: 09/896,580
	Applicant(s): Eric T. BALDWIN et al.	Confirmation No.: 7868
	Filing Date: 29 June 2001	Group: 2621



## U.S. PATENT DOCUMENTS

Examiner Initial	Document Number	Date	Name	Class	Subclass	Filing Date If Appropriate
	NONE					RECEIVED
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## FOREIGN PATENT DOCUMENTS

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Examiner Initial	Document Number	Date	Country	Class	Subclass	Translation
NE	0 879 879 A2	11/25/98	EPO			Yes No

## OTHER DOCUMENTS (Including Authors, Title, Date, Pertinent Papers, etc.)

Examiner Initial	Document Description
NE	Becker et al., "Structure of Peptide Deformylase and Identification of the Substrate Binding Site," <i>The Journal of Biological Chemistry</i> , 273(19):11413-11416 (1998).
NE	Becker et al., "Iron center, substrate recognition and mechanism of peptide deformylase," <i>Nature Structural Biology</i> , 5(12):1053-1058 (1998).
NE	Brizzard et al., "Immunoaffinity Purification of FLAG® Epitope-Tagged Bacterial Alkaline Phosphatase Using a Novel Monoclonal Antibody and Peptide Elution," <i>BioTechniques</i> , 16(4):730-735 (1994).
NE	Chang et al., "Methionine Aminopeptidase Gene of <i>Escherichia coli</i> Is Essential for Cell Growth," <i>Journal of Bacteriology</i> , 171(7):4071-4072 (1989).
NE	Chen et al., "Mechanistic Studies on the Aminopeptidase from <i>Aeromonas proteolytica</i> : A Two-Metal Ion Mechanism for Peptide Hydrolysis," <i>Biochemistry</i> , 36(14):4278-4286 (1997).
NE	Chiang et al., "Expression and Purification of General Transcription Factors by FLAG Epitope-Tagging and Peptide Elution," <i>Peptide Research</i> , 6(2):62-64 (1993).
NE	Dardel et al., "Solution Structure of Nickel-peptide Deformylase," <i>Journal of Molecular Biology</i> , 280(3):501-513 (1998).
NE	Ford et al., "Fusion Tails for the Recovery and Purification of Recombinant Proteins," <i>Protein Expression and Purification</i> , 2:95-107 (1991).

EXAMINER <i>N. Ahmed</i>	Date Considered 9/9/03
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\*Examiner: Initial if citation considered, whether or not citation is in conformance with MPEP 609; Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.

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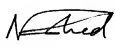
Examiner Initial	Document Description
	Groche et al., "Isolation and Crystallization of Functionally Competent <i>Escherichia coli</i> Peptide Deformylase Forms Containing either Iron or Nickel in the Active Site," <i>Biochemical and Biophysical Research Communications</i> , 246(2):342-346 (1998).
ne	Hopp et al., "A Short Polypeptide Marker Sequence Useful for Recombinant Protein Identification and Purification," <i>Biotechnology</i> , 6(10):1204-1210 (1988).
ne	Hu et al., "H-Phosphonate Derivatives as Novel Peptide Deformylase Inhibitors," <i>Bioorganic &amp; Medicinal Chemistry Letters</i> , 8:2479-2482 (1998).
ne	Hu et al., "Determination of Substrate Specificity for Peptide Deformylase through the Screening of a Combinatorial Peptide Library," <i>Biochemistry</i> , 38(2):643-650 (1999).
ne	Laemmli, "Cleavage of Structural Proteins during the Assembly of the Head of Bacteriophage T4," <i>Nature</i> , 227(5259):680-685 (1970).
ne	Lazennec et al., "Formate Dehydrogenase-Coupled Spectrophotometric Assay of Peptide Deformylase," <i>Analytical Biochemistry</i> , 244:180-182 (1997).
ne	Meinzel et al., "Mapping of the Active Site Zinc Ligands of Peptide Deformylase," <i>Journal of Molecular Biology</i> , 254(2):175-183 (1995).
ne	Meinzel et al., "A New Subclass of the Zinc Metalloproteases Superfamily Revealed by the Solution Structure of Peptide Deformylase," <i>Journal of Molecular Biology</i> , 262(3):375-386 (1996).
ne	Meinzel et al., "Structure-Function Relationships within the Peptide Deformylase Family. Evidence for a Conserved Architecture of the Active Site Involving Three Conserved Motifs and a Metal Ion," <i>Journal of Molecular Biology</i> , 267(3):749-761 (1997).
ne	Meinzel et al., "Design and Synthesis of Substrate Analogue Inhibitors of Peptide Deformylase," <i>Biochemistry</i> , 38(14):4287-4295 (1999).
ne	Prescott et al., " <i>Aeromonas</i> Aminopeptidase," <i>Methods in Enzymology</i> , 45(Part B):530-543 (1976).
ne	QIAexpress® - The Complete System Ni-NTA Technology and the 6xHis Tag. Datasheet [online]. Qiagen [retrieved on 2001-11-06]. Retrieved from the Internet: <URL:www.qiagen.com/catalog/chapter_03/chap3.asp>, 3 pages.

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Examination Initials	Document Description
 ME	QIAexpress® Expression System. Datasheet [online]. Qiagen [retrieved 2001-11-06]. Retrieved from the Internet: <URL:www.qiagen.com/catalog/chapter_03/chap3.asp>, 5 pages.
ME	QIAexpress® Protein Purification System. Datasheet [online]. Qiagen [retrieved on 2001-11-06]. Retrieved from the Internet: <URL:www.qiagen.com/catalog/chapter_03/chap3.asp>, 5 pages.
ME	Qiagen, <i>QIAexpress Detection and Assay Handbook</i> , pages 9-45, 52-76 (1999).
ME	Wei et al., "Continuous Spectrophotometric Assay of Peptide Deformylase," <i>Analytical Biochemistry</i> , 250:29-34 (1997).

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